☑004/016

Application No.:10/597,813   Docket No.: JCLA21512-R

AMENDMENT

In The Specification:

Please amend the following paragraphs.

Page 13, first paragraph to Page 14, first paragraph

Preferred are the following elongated sequences of SEQ ID NO 3 which can be represented by

the following general formula:

5'-XCAGCCCCCGACCCATGZ-3' <u>SEQ ID NO:101</u>

wherein X is selected from the group comprising the following oligonucleotides:

ACAGGACGATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:102</u>

CAGGACGATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:103</u>

AGGACGATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:104</u>

GGACGATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:105</u>

GACGATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:106</u>

ACGATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:107</u>

CGATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:108</u>

GATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:109</u>

ATGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:110</u>

TGTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:111</u>

GTGCAGCGGCCACAGGCCCCTGAG, <u>SEQ ID NO:112</u>

TGCAGCGGCCACAGGCCCCTGAG, SEQ ID NO:113

GCAGCGGCCACAGGCCCCTGAG, SEQ ID NO:114

CAGCGGCCACAGGCCCCTGAG, SEQ ID NO:115

AGCGGCCACAGGCCCCTGAG, SEQ ID NO:116

GCGGCCACAGGCCCCTGAG, SEQ ID NO:117

CGGCCACAGGCCCCTGAG, SEQ ID NO:118

GGCCACAGGCCCCTGAG, SEQ ID NO:119

GCCACAGGCCCCTGAG, SEQ ID NO:120

CCACAGGCCCCTGAG, SEQ ID NO:121

CACAGGCCCCTGAG, SEQ ID NO:122

ACAGGCCCCTGAG, SEQ ID NO:123

CAGGCCCCTGAG, SEQ ID NO:124

AGGCCCCTGAG, SEQ ID NO:125

GGCCCCTGAG, SEQ ID NO:126

GCCCCTGAG, SEQ ID NO:127

CCCCTGAG, SEQ ID NO:128

CCCTGAG, SEQ ID NO:129

CCTGAG, SEQ ID NO:130

CTGAG, SEQ ID NO:131

TGAG, SEQ ID NO:132

GAG, AG, G,

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and wherein Z is selected from the group comprising the following oligonucleotides:

GCAGACCCCGCTGCTCGTCATAGACCGAGCCCCC, SEQ ID NO:133

GCAGACCCCGCTGCTCGTCATAGACCGAGCCCC, SEQ ID NO:134

GCAGACCCCGCTGCTCGTCATAGACCGAGCCC, SEQ ID NO:135

GCAGACCCCGCTGCTCGTCATAGACCGAGCC, SEQ ID NO:136

GCAGACCCCGCTGCTCGTCATAGACCGAGC, SEQ ID NO:137

GCAGACCCCGCTGCTCGTCATAGACCGAG, SEQ ID NO:138

GCAGACCCCGCTGCTCGTCATAGACCGA, SEQ ID NO:139

GCAGACCCCGCTGCTCGTCATAGACCG, SEQ ID NO:140

GCAGACCCCGCTGCTCGTCATAGACC, SEQ ID NO:141

GCAGACCCCGCTGCTCGTCATAGAC, SEQ ID NO:142

GCAGACCCCGCTGCTCGTCATAGA, SEQ ID NO:143

GCAGACCCCGCTGCTCGTCATAG, SEQ ID NO:144

GCAGACCCCGCTGCTCGTCATA, SEQ ID NO:145

GCAGACCCCGCTGCTCGTCAT, SEQ ID NO:146

GCAGACCCCGCTGCTCGTCA, SEQ ID NO:147

GCAGACCCCGCTGCTCGTC, SEQ ID NO:148

GCAGACCCCGCTGCTCGT, SEQ ID NO:149

GCAGACCCCGCTGCTCG, SEQ ID NO:150

GCAGACCCCGCTGCTC, SEQ ID NO:151

GCAGACCCCGCTGCT, SEQ ID NO:152

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GCAGACCCCGCTGC, <u>SEQ ID NO:153</u>

GCAGACCCCGCTG, <u>SEQ ID NO:154</u>

GCAGACCCCGCT, <u>SEQ ID NO:155</u>

GCAGACCCCGC, <u>SEQ ID NO:156</u>

GCAGACCCCG, <u>SEQ ID NO:157</u>

GCAGACCCC, <u>SEQ ID NO:158</u>

GCAGACCC, <u>SEQ ID NO:159</u>

GCAGACC, <u>SEQ ID NO:160</u>

GCAGAC, <u>SEQ ID NO:161</u>

GCAGA, <u>SEQ ID NO:162</u>

GCAG, <u>SEQ ID NO:163</u>

GCA, GC, G,

and wherein X and Z together comprise not more than 34 nucleobases and mimetics thereof.